

**Amendments to the Specification:**

Please amend the specification as follows:

Please amend the paragraph on page 15, lines 5 and 6, as follows:

[Figure 22] Figure 22 is a schematic diagram showing the preparation of humanized MABL-2 HL5s containing S-S bonds **(GGGGS linker peptide disclosed as SEQ ID NO: 109).**

Please amend the paragraph beginning on page 33, line 27, and ending on page 34, line 22, as follows:

In the present invention, suitable linkers joining the H chain V region and the L chain V region or linkers joining diabody-forming fragments to form single-chain diabodies include any peptide linkers that can be introduced by genetic engineering or synthetic linkers, such as linkers disclosed in Protein Engineering, 9(3), 299-305, 1996. For example, peptide linkers include:

Ser

Gly·Ser

Gly·Gly·Ser

Ser·Gly·Gly

Gly·Gly·Gly·Ser **(SEQ ID NO: 115)**

Ser·Gly·Gly·Gly **(SEQ ID NO: 116)**

Gly·Gly·Gly·Gly·Ser **(SEQ ID NO: 109)**

Ser·Gly·Gly·Gly·Gly **(SEQ ID NO: 117)**

Gly·Gly·Gly·Gly·Gly·Ser **(SEQ ID NO: 118)**

Ser·Gly·Gly·Gly·Gly·Gly **(SEQ ID NO: 119)**

Gly·Gly·Gly·Gly·Gly·Gly·Ser **(SEQ ID NO: 120)**

Ser·Gly·Gly·Gly·Gly·Gly·Gly **(SEQ ID NO: 121)**

(Gly·Gly·Gly·Gly·Ser)<sub>n</sub> (SEQ ID NO: 109)

(Ser·Gly·Gly·Gly·Gly)<sub>n</sub> (SEQ ID NO: 117)

wherein n is an integer of 1 or more. The length of linker peptides can be selected as appropriate by those skilled in the art depending on the purpose.

Please amend the paragraph on page 64, lines 5-15, as follows:

The forward primer X5-huLgS (primer C, SEQ ID NO: 70) for the L chain V region was designed to hybridize to the DNA encoding the C-terminus of the H chain V region, to contain the DNA sequence encoding the 5mer linker region consisting of Gly Gly Gly Gly Ser (SEQ ID NO: ~~72~~ 109) and to overlap the DNA encoding the N-terminus of the L chain V region. The reverse primer NothuLAS (primer D, SEQ ID NO: 71) for the L chain V region was designed to hybridize to the DNA encoding the C-terminus of the L chain V region and to have two transcription termination codons and an NotI restriction endonuclease site.

Please amend the paragraph on page 67, lines 4-13, as follows:

The forward primer X15huHS (primer E, SEQ ID NO: 75) for the H chain V region was designed to overlap a part of the 15mer linker described below and to hybridize to the DNA encoding the N-terminus of the H chain V region. The reverse primer X15huLAS (primer F, SEQ ID NO: 76) for the L chain V region was designed to hybridize to the DNA encoding the C-terminus of the L chain V region and to hybridize to the DNA sequence encoding the 15mer linker region consisting of Gly Gly Gly Gly Ser Gly Gly Gly Gly Ser Gly Gly Gly Gly Ser (SEQ ID NO: ~~77~~ 111).